

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 09/954,737C
Source: IFw/6
Date Processed by STIC: 6/1/05

ENTERED



I FW16

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/954,737C

DATE: 06/01/2005
TIME: 14:29:45

Input Set : A:\GC634-2-SEQLIST2.TXT
Output Set: N:\CRF4\06012005\I954737C.raw

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4 <110> APPLICANT: Bron, Sierd
5     Jongbloed, Jan D.H.
6     Mueller, Joerg P.
7     Van Dijl, Jan M.
9 <120> TITLE OF INVENTION: Twin-Arginine Translocation in Bacillus
12 <130> FILE REFERENCE: GC634-2
14 <140> CURRENT APPLICATION NUMBER: US 09/954,737C
15 <141> CURRENT FILING DATE: 2001-09-17
17 <150> PRIOR APPLICATION NUMBER: US 60/233,610
18 <151> PRIOR FILING DATE: 2000-09-18
20 <160> NUMBER OF SEQ ID NOS: 83
22 <170> SOFTWARE: FastSEQ for Windows Version 4.0
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 89
26 <212> TYPE: PRT
27 <213> ORGANISM: Escherichia coli
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31    1           5           10          15
32 Val Leu Leu Phe Gly Thr Lys Lys Leu Gly Ser Ile Gly Ser Asp Leu
33    20          25          30
34 Gly Ala Ser Ile Lys Gly Phe Lys Lys Ala Met Ser Asp Asp Glu Pro
35    35          40          45
36 Lys Gln Asp Lys Thr Ser Gln Asp Ala Asp Phe Thr Ala Lys Thr Ile
37    50          55          60
38 Ala Asp Lys Gln Ala Asp Thr Asn Gln Glu Gln Ala Lys Thr Glu Asp
39    65          70          75          80
40 Ala Lys Arg His Asp Lys Glu Gln Val
41    85
43 <210> SEQ ID NO: 2
44 <211> LENGTH: 67
45 <212> TYPE: PRT
46 <213> ORGANISM: Escherichia coli
48 <400> SEQUENCE: 2
49 Met Gly Glu Ile Ser Ile Thr Lys Leu Leu Val Val Ala Ala Leu Val
50    1           5           10          15
51 Val Leu Leu Phe Gly Thr Lys Lys Leu Arg Thr Leu Gly Gly Asp Leu
52    20          25          30
53 Gly Ala Ala Ile Lys Gly Phe Lys Lys Ala Met Asn Asp Asp Ala
54    35          40          45
55 Ala Ala Lys Lys Gly Ala Asp Val Asp Leu Gln Ala Glu Lys Leu Ser
56    50          55          60
57 His Lys Glu

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58 65
60 <210> SEQ ID NO: 3
61 <211> LENGTH: 57
62 <212> TYPE: PRT
63 <213> ORGANISM: Bacillus subtilis
65 <400> SEQUENCE: 3
66 Met Pro Ile Gly Pro Gly Ser Leu Ala Val Ile Ala Ile Val Ala Leu
67 1 5 10 15
68 Ile Ile Phe Gly Pro Lys Lys Leu Pro Glu Leu Gly Lys Ala Ala Gly
69 20 25 30
70 Asp Thr Leu Arg Glu Phe Lys Asn Ala Thr Lys Gly Leu Thr Ser Asp
71 35 40 45
72 Glu Glu Glu Lys Lys Glu Asp Gln
73 50 55
75 <210> SEQ ID NO: 4
76 <211> LENGTH: 70
77 <212> TYPE: PRT
78 <213> ORGANISM: Bacillus subtilis
80 <400> SEQUENCE: 4
81 Met Phe Ser Asn Ile Gly Ile Pro Gly Leu Ile Leu Ile Phe Val Ile
82 1 5 10 15
83 Ala Ile Ile Ile Phe Gly Pro Ser Lys Leu Pro Glu Ile Gly Arg Ala
84 20 25 30
85 Ala Lys Arg Thr Leu Leu Glu Phe Lys Ser Ala Thr Lys Ser Leu Val
86 35 40 45
87 Ser Gly Asp Glu Lys Glu Lys Ser Ala Glu Leu Thr Ala Val Lys
88 50 55 60
89 Gln Asp Lys Asn Ala Gly
90 65 70
92 <210> SEQ ID NO: 5
93 <211> LENGTH: 62
94 <212> TYPE: PRT
95 <213> ORGANISM: Bacillus subtilis
97 <400> SEQUENCE: 5
98 Met Glu Leu Ser Phe Thr Lys Ile Leu Val Ile Leu Phe Val Gly Phe
99 1 5 10 15
100 Leu Val Phe Gly Pro Asp Lys Leu Pro Ala Leu Gly Arg Ala Ala Gly
101 20 25 30
102 Lys Ala Leu Ser Glu Phe Lys Gln Ala Thr Ser Gly Leu Thr Gln Asp
103 35 40 45
104 Ile Arg Lys Asn Asp Ser Glu Asn Lys Glu Asp Lys Gln Met
105 50 55 60
107 <210> SEQ ID NO: 6
108 <211> LENGTH: 171
109 <212> TYPE: PRT
110 <213> ORGANISM: Escherichia coli
112 <400> SEQUENCE: 6
113 Met Phe Asp Ile Gly Phe Ser Glu Leu Leu Val Phe Ile Ile Gly
114 1 5 10 15

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115 Leu Val Val Leu Gly Pro Gln Arg Leu Pro Val Ala Val Lys Thr Val
116          20           25           30
117 Ala Gly Trp Ile Arg Ala Leu Arg Ser Leu Ala Thr Thr Val Gln Asn
118          35           40           45
119 Glu Leu Thr Gln Glu Leu Lys Leu Gln Glu Phe Gln Asp Ser Leu Lys
120          50           55           60
121 Lys Val Glu Lys Ala Ser Leu Thr Asn Leu Thr Pro Glu Leu Lys Ala
122          65           70           75           80
123 Ser Met Asp Glu Leu Arg Gln Ala Ala Glu Ser Met Lys Arg Ser Tyr
124          85           90           95
125 Val Ala Asn Asp Pro Glu Lys Ala Ser Asp Glu Ala His Thr Ile His
126          100          105          110
127 Asn Pro Val Val Lys Asp Asn Glu Ala Ala His Glu Gly Val Thr Pro
128          115          120          125
129 Ala Ala Ala Gln Thr Gln Ala Ser Ser Pro Glu Gln Lys Pro Glu Thr
130          130          135          140
131 Thr Pro Glu Pro Val Val Lys Pro Ala Ala Asp Ala Glu Pro Lys Thr
132          145          150          155          160
133 Ala Ala Pro Ser Pro Ser Ser Asp Lys Pro
134          165          170
136 <210> SEQ ID NO: 7
137 <211> LENGTH: 258
138 <212> TYPE: PRT
139 <213> ORGANISM: Escherichia coli
141 <400> SEQUENCE: 7
142 Met Ser Val Glu Asp Thr Gln Pro Leu Ile Thr His Leu Ile Glu Leu
143 1          5           10           15
144 Arg Lys Arg Leu Leu Asn Cys Ile Ile Ala Val Ile Val Ile Phe Leu
145          20           25           30
146 Cys Leu Val Tyr Phe Ala Asn Asp Ile Tyr His Leu Val Ser Ala Pro
147          35           40           45
148 Leu Ile Lys Gln Leu Pro Gln Gly Ser Thr Met Ile Ala Thr Asp Val
149          50           55           60
150 Ala Ser Pro Phe Phe Thr Pro Ile Lys Leu Thr Phe Met Val Ser Leu
151          65           70           75           80
152 Ile Leu Ser Ala Pro Val Ile Leu Tyr Gln Val Trp Ala Phe Ile Ala
153          85           90           95
154 Pro Ala Leu Tyr Lys His Glu Arg Arg Leu Val Val Pro Leu Leu Val
155          100          105          110
156 Ser Ser Ser Leu Leu Phe Tyr Ile Gly Met Ala Phe Ala Tyr Phe Val
157          115          120          125
158 Val Phe Pro Leu Ala Phe Gly Phe Leu Ala Asn Thr Ala Pro Glu Gly
159          130          135          140
160 Val Gln Val Ser Thr Asp Ile Ala Ser Tyr Leu Ser Phe Val Met Ala
161          145          150          155          160
162 Leu Phe Met Ala Phe Gly Val Ser Phe Glu Val Pro Val Ala Ile Val
163          165          170          175
164 Leu Leu Cys Trp Met Gly Ile Thr Ser Pro Glu Asp Leu Arg Lys Lys
165          180          185          190

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166 Arg Pro Tyr Val Leu Val Gly Ala Phe Val Val Gly Met Leu Leu Thr
 167 195 200 205
 168 Pro Pro Asp Val Phe Ser Gln Thr Leu Leu Ala Ile Pro Met Tyr Cys
 169 210 215 220
 170 Leu Phe Glu Ile Gly Val Phe Phe Ser Arg Phe Tyr Val Gly Lys Gly
 171 225 230 235 240
 172 Arg Asn Arg Glu Glu Asn Asp Ala Glu Ala Glu Ser Glu Lys Thr
 173 245 250 255
 174 Glu Glu
 177 <210> SEQ ID NO: 8
 178 <211> LENGTH: 254
 179 <212> TYPE: PRT
 180 <213> ORGANISM: Bacillus subtilis
 182 <400> SEQUENCE: 8
 183 Met Thr Arg Met Lys Val Asn Gln Met Ser Leu Leu Glu His Ile Ala
 184 1 5 10 15
 185 Glu Leu Arg Lys Arg Leu Leu Ile Val Ala Leu Ala Phe Val Val Phe
 186 20 25 30
 187 Phe Ile Ala Gly Phe Phe Leu Ala Lys Pro Ile Ile Val Tyr Leu Gln
 188 35 40 45
 189 Glu Thr Asp Glu Ala Lys Gln Leu Thr Leu Asn Ala Phe Asn Leu Thr
 190 50 55 60
 191 Asp Pro Leu Tyr Val Phe Met Gln Phe Ala Phe Ile Ile Gly Ile Val
 192 65 70 75 80
 193 Leu Thr Ser Pro Val Ile Leu Tyr Gln Leu Trp Ala Phe Val Ser Pro
 194 85 90 95
 195 Gly Leu Tyr Glu Lys Glu Arg Lys Val Thr Leu Ser Tyr Ile Pro Val
 196 100 105 110
 197 Ser Ile Leu Leu Phe Leu Ala Gly Leu Ser Phe Ser Tyr Tyr Ile Leu
 198 115 120 125
 199 Phe Pro Phe Val Val Asp Phe Met Lys Arg Ile Ser Gln Asp Leu Asn
 200 130 135 140
 201 Val Asn Gln Val Ile Gly Ile Asn Glu Tyr Phe His Phe Leu Leu Gln
 202 145 150 155 160
 203 Leu Thr Ile Pro Phe Gly Leu Leu Phe Gln Met Pro Val Ile Leu Met
 204 165 170 175
 205 Phe Leu Thr Arg Leu Gly Ile Val Thr Pro Met Phe Leu Ala Lys Ile
 206 180 185 190
 207 Arg Lys Tyr Ala Tyr Phe Thr Leu Leu Val Ile Ala Ala Leu Ile Thr
 208 195 200 205
 209 Pro Pro Glu Leu Leu Ser His Met Met Val Thr Val Pro Leu Leu Ile
 210 210 215 220
 211 Leu Tyr Glu Ile Ser Ile Leu Ile Ser Lys Ala Ala Tyr Arg Lys Ala
 212 225 230 235 240
 213 Gln Lys Ser Ser Ala Ala Asp Arg Asp Val Ser Ser Gly Gln
 214 245 250
 216 <210> SEQ ID NO: 9
 217 <211> LENGTH: 245
 218 <212> TYPE: PRT

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Input Set : A:\GC634-2-SEQLIST2.TXT

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219 <213> ORGANISM: Bacillus subtilis
 221 <400> SEQUENCE: 9
 222 Met Asp Lys Lys Glu Thr His Leu Ile Gly His Leu Glu Glu Leu Arg
 223 1 5 10 15
 224 Arg Arg Ile Ile Val Thr Leu Ala Ala Phe Phe Leu Phe Leu Ile Thr
 225 20 25 30
 226 Ala Phe Leu Phe Val Gln Asp Ile Tyr Asp Trp Leu Ile Arg Asp Leu
 227 35 40 45
 228 Asp Gly Lys Leu Ala Val Leu Gly Pro Ser Glu Ile Leu Trp Val Tyr
 229 50 55 60
 230 Met Met Leu Ser Gly Ile Cys Ala Ile Ala Ala Ser Ile Pro Val Ala
 231 65 70 75 80
 232 Ala Tyr Gln Leu Trp Arg Phe Val Ala Pro Ala Leu Thr Lys Thr Glu
 233 85 90 95
 234 Arg Lys Val Thr Ile Met Tyr Ile Met Tyr Ile Pro Gly Leu Phe Ala
 235 100 105 110
 236 Leu Phe Leu Ala Gly Ile Ser Phe Gly Tyr Phe Val Leu Phe Pro Ile
 237 115 120 125
 238 Val Leu Ser Phe Leu Thr His Leu Ser Ser Gly His Phe Glu Thr Met
 239 130 135 140
 240 Phe Thr Ala Asp Arg Tyr Phe Arg Phe Met Val Asn Leu Ser Leu Pro
 241 145 150 155 160
 242 Phe Gly Phe Leu Phe Glu Met Pro Leu Val Val Met Phe Leu Thr Arg
 243 165 170 175
 244 Leu Gly Ile Leu Asn Pro Tyr Arg Leu Ala Lys Ala Arg Lys Leu Ser
 245 180 185 190
 246 Tyr Phe Leu Leu Ile Val Val Ser Ile Leu Ile Thr Pro Pro Asp Phe
 247 195 200 205
 248 Ile Ser Asp Phe Leu Val Met Ile Pro Leu Leu Val Leu Phe Glu Val
 249 210 215 220
 250 Ser Val Thr Leu Ser Ala Phe Val Tyr Lys Lys Arg Met Arg Glu Glu
 251 225 230 235 240
 252 Thr Ala Ala Ala Ala
 253 245
 255 <210> SEQ ID NO: 10
 256 <211> LENGTH: 63
 257 <212> TYPE: PRT
 258 <213> ORGANISM: Bacillus alcalophilus
 260 <400> SEQUENCE: 10
 261 Met Gly Gly Leu Ser Val Gly Ser Val Val Leu Ile Ala Leu Val Ala
 262 1 5 10 15
 263 Leu Leu Ile Phe Gly Pro Lys Lys Leu Pro Glu Leu Gly Lys Ala Ala
 264 20 25 30
 265 Gly Ser Thr Leu Arg Glu Phe Lys Asn Ala Thr Lys Gly Leu Ala Asp
 266 35 40 45
 267 Asp Asp Asp Asp Thr Lys Ser Thr Asn Val Gln Lys Glu Lys Ala
 268 50 55 60
 270 <210> SEQ ID NO: 11
 271 <211> LENGTH: 272

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/954,737C

DATE: 06/01/2005

TIME: 14:29:46

Input Set : A:\GC634-2-SEQLIST2.TXT

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